
Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2008; month=12; day=5; hr=9; min=4; sec=21; ms=55;]

Reviewer Comments:

<210> 1

<211> 390

<212> PRT

<213> Homo sapien

* * * * * * * * *

Please change the spelling of "Homo sapien" to "Homo sapiens."

<210> 16

<211> 20

<212> PRT

<213> synthetic

* * * * * * * * *

For SEQ ID # 16, numeric identifier <213> can only be one of three choices, "Scientific name, i.e. Genus/species, Unknown or Artificial Sequence." For all sequences using "Unknown or Artificial sequence", for numeric identifier <213>, a mandatory feature is required to explain the source of the genetic material. The feature consists of <220>, which remains blank, and <223>, which states the source of the genetic material. Suggest using "Artificial sequence" for numeric identifier <213> and "Synthetic" for numeric identifier <223> in the mandatory feature. Please make all necessary changes.

Validated By CRFValidator v 1.0.3

Application No: 10585499 Version No: 1.0

Input Set:

Output Set:

Started: 2008-11-18 12:10:27.981 **Finished:** 2008-11-18 12:10:29.193

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 212 ms

Total Warnings: 16
Total Errors: 1

No. of SeqIDs Defined: 17

Actual SeqID Count: 17

Error code		Error Descrip	otion							
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(1)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(2)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(3)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(4)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(5)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(6)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(7)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(8)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(9)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(10)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(11)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(12)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(13)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(14)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(15)
Ε	356	Organism :	is not per	rmitte	d in	n <213>	iı	n SEÇ	Q II	(16)
W	402	Undefined	organism	found	in	<213>	in	SEQ	ID	(17)

SEQUENCE LISTING

OF

<110>	The Government of the United States as represented by the Secretary, Department of Health and Human Services													
<120>	COMPOSITIONS AND METHODS FOR THE HIGH EFFICIENCY EXPRESSION THE TRANSFORMING GROWTH FACTOR-BETA SUPERGENE FAMILY													
<130>	NIHA-0186													
	10585499 2008-11-18													
<150> <151>	US 60/534,379 2004-01-06													
<160>	17													
<170>	PatentIn version 3.3													
<212>	1 390 PRT Homo sa	apien												
<400>	<400> 1													
Met Pr	o Pro Se	er Gly I 5	eu Arg	Leu	Leu	Leu 10	Leu	Leu	Leu	Pro	Leu 15	Leu		
Trp Le	u Leu Va 20		hr Pro	Gly	Arg 25	Pro	Ala	Ala	Gly	Leu 30	Ser	Thr		
Cys Ly	s Thr Il 35	le Asp M	Met Glu	Leu 40	Val	Lys	Arg	Lys	Arg 45	Ile	Glu	Ala		
Ile Ar	g Gly Gl	ln Ile I	eu Ser 55	Lys	Leu	Arg	Leu	Ala 60	Ser	Pro	Pro	Ser		
Gln Gl	y Glu Va		ro Gly 0	Pro	Leu	Pro	Glu 75	Ala	Val	Leu	Ala	Leu 80		
Tyr As:	n Ser Th	nr Arg A 85	.sp Arg	Val	Ala	Gly 90	Glu	Ser	Ala	Glu	Pro 95	Glu		
Pro Gl	u Pro Gl 10		sp Tyr	Tyr	Ala 105	Lys	Glu	Val	Thr	Arg 110	Val	Leu		

Met Val Glu Thr His Asn Glu Ile Tyr Asp Lys Phe Lys Gln Ser Thr

125

120

His	Ser 130	Ile	Tyr	Met	Phe	Phe 135	Asn	Thr	Ser	Glu	Leu 140	Arg	Glu	Ala	Val
Pro 145	Glu	Pro	Val	Leu	Leu 150	Ser	Arg	Ala	Glu	Leu 155	Arg	Leu	Leu	Arg	Leu 160
Lys	Leu	Lys	Val	Glu 165	Gln	His	Val	Glu	Leu 170	Tyr	Gln	Lys	Tyr	Ser 175	Asn
Asn	Ser	Trp	Arg 180	Tyr	Leu	Ser	Asn	Arg 185	Leu	Leu	Ala	Pro	Ser 190	Asp	Ser
Pro	Glu	Trp 195	Leu	Ser	Phe	Asp	Val 200	Thr	Gly	Val	Val	Arg 205	Gln	Trp	Leu
Ser	Arg 210	Gly	Gly	Glu	Ile	Glu 215	Gly	Phe	Arg	Leu	Ser 220	Ala	His	Суз	Ser
Cys 225	Asp	Ser	Arg	Asp	Asn 230	Thr	Leu	Gln	Val	Asp 235	Ile	Asn	Gly	Phe	Thr 240
Thr	Gly	Arg	Arg	Gly 245	Asp	Leu	Ala	Thr	Ile 250	His	Gly	Met	Asn	Arg 255	Pro
Phe	Leu	Leu	Leu 260	Met	Ala	Thr	Pro	Leu 265	Glu	Arg	Ala	Gln	His 270	Leu	Gln
Ser	Ser	Arg 275	His	Arg	Arg	Ala	Leu 280	Asp	Thr	Asn	Tyr	Cys 285	Phe	Ser	Ser
Thr	Glu 290	Lys	Asn	Cys	Cys	Val 295	Arg	Gln	Leu	Tyr	Ile 300	Asp	Phe	Arg	Lys
Asp 305	Leu	Gly	Trp	Lys	Trp 310	Ile	His	Glu	Pro	Lys 315	Gly	Tyr	His	Ala	Asn 320
Phe	Cys	Leu	Gly	Pro 325	Cys	Pro	Tyr	Ile	Trp 330	Ser	Leu	Asp	Thr	Gln 335	Tyr
Ser	Lys	Val	Leu 340	Ala	Leu	Tyr	Asn	Gln 345	His	Asn	Pro	Gly	Ala 350	Ser	Ala

Ala Pro Cys Cys Val Pro Gln Ala Leu Glu Pro Leu Pro Ile Val Tyr 355

Tyr Val Gly Arg Lys Pro Lys Val Glu Gln Leu Ser Asn Met Ile Val 370

370

380

41a Leu Glu Pro Leu Pro Leu Pro Ile Val Glu Glu Gln Leu Ser Asn Met Ile Val 370

380

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

48

<210> 2 <211> 390 <212> PRT <213> pig

<400> 2

Met Pro Pro Ser Gly Leu Arg Leu Leu Pro Leu Leu Pro Leu Leu 1 5 10 15

Trp Leu Leu Val Leu Thr Pro Gly Arg Pro Ala Ala Gly Leu Ser Thr 20 25 30

Cys Lys Thr Ile Asp Met Glu Leu Val Lys Arg Lys Arg Ile Glu Ala 35 40 45

Ile Arg Gly Gln Ile Leu Ser Lys Leu Arg Leu Ala Ser Pro Pro Ser 50 55 60

Gln Gly Asp Val Pro Pro Gly Pro Leu Pro Glu Ala Val Leu Ala Leu 65 70 75 80

Tyr Asn Ser Thr Arg Asp Arg Val Ala Gly Glu Ser Val Glu Pro Glu 85 90 95

Pro Glu Pro Glu Ala Asp Tyr Tyr Ala Lys Glu Val Thr Arg Val Leu 100 105 110

Met Leu Glu Ser Gly Asn Gln Ile Tyr Asp Lys Phe Lys Gly Thr Pro 115 120 125

His Ser Leu Tyr Met Leu Phe Asn Thr Ser Glu Leu Arg Glu Ala Val 130 135 140

Pro Glu Pro Val Leu Leu Ser Arg Ala Glu Leu Arg Leu Arg Leu

145 150 155 16	50
----------------	----

Lys Leu Lys Val Glu Gln His Val Glu Leu Tyr Gln Lys Tyr Ser Asn 165 170 175

Asp Ser Trp Arg Tyr Leu Ser Asn Arg Leu Leu Ala Pro Ser Asp Ser 180 185 190

Pro Glu Trp Leu Ser Phe Asp Val Thr Gly Val Val Arg Gln Trp Leu 195 200 205

Thr Arg Arg Glu Ala Ile Glu Gly Phe Arg Leu Ser Ala His Cys Ser 210 215 220

Cys Asp Ser Lys Asp Asn Thr Leu His Val Glu Ile Asn Gly Phe Asn 225 230 235 240

Ser Gly Arg Arg Gly Asp Leu Ala Thr Ile His Gly Met Asn Arg Pro 245 250 255

Phe Leu Leu Met Ala Thr Pro Leu Glu Arg Ala Gln His Leu His
260 265 270

Ser Ser Arg His Arg Arg Ala Leu Asp Thr Asn Tyr Cys Phe Ser Ser 275 280 285

Thr Glu Lys Asn Cys Cys Val Arg Gln Leu Tyr Ile Asp Phe Arg Lys 290 295 300

Asp Leu Gly Trp Lys Trp Ile His Glu Pro Lys Gly Tyr His Ala Asn 305 310 315 320

Phe Cys Leu Gly Pro Cys Pro Tyr Ile Trp Ser Leu Asp Thr Gln Tyr 325 330 335

Ser Lys Val Leu Ala Leu Tyr Asn Gln His Asn Pro Gly Ala Ser Ala 340 345 350

Ala Pro Cys Cys Val Pro Gln Ala Leu Glu Pro Leu Pro Ile Val Tyr 355 360 365

Tyr Val Gly Arg Lys Pro Lys Val Glu Gln Leu Ser Asn Met Ile Val 370 375 380

Arg Ser Cys Lys Cys Ser 385 390 <210> 3 <211> 414 <212> PRT <213> Homo sapien <400> 3 Met His Tyr Cys Val Leu Ser Ala Phe Leu Ile Leu His Leu Val Thr 10 15 5 Val Ala Leu Ser Leu Ser Thr Cys Ser Thr Leu Asp Met Asp Gln Phe 20 25 30 Met Arg Lys Arg Ile Glu Ala Ile Arg Gly Gln Ile Leu Ser Lys Leu 35 40 45 Lys Leu Thr Ser Pro Pro Glu Asp Tyr Pro Glu Pro Glu Glu Val Pro 50 55 60 Pro Glu Val Ile Ser Ile Tyr Asn Ser Thr Arg Asp Leu Leu Gln Glu
 65
 70
 75
 80
 Lys Ala Ser Arg Arg Ala Ala Cys Glu Arg Glu Arg Ser Asp Glu Glu Tyr Tyr Ala Lys Glu Val Tyr Lys Ile Asp Met Pro Pro Phe Phe 100 105 110 Pro Ser Glu Asn Ala Ile Pro Pro Thr Phe Tyr Arg Pro Tyr Phe Arg 115 120 125 Ile Val Arg Phe Asp Val Ser Ala Met Glu Lys Asn Ala Ser Asn Leu 135 140 130 Val Lys Ala Glu Phe Arg Val Phe Arg Leu Gln Asn Pro Lys Ala Arg 145 150 155 160

Val Pro Glu Gln Arg Ile Glu Leu Tyr Gln Ile Leu Lys Ser Lys Asp

165 170 175

Leu	Thr	Ser	Pro 180	Thr	Gln	Arg	Tyr	Ile 185	Asp	Ser	Lys	Val	Val 190	Lys	Thr
Arg	Ala	Glu 195	Gly	Glu	Trp	Leu	Ser 200	Phe	Asp	Val	Thr	Asp 205	Ala	Val	His
Glu	Trp 210	Leu	His	His	Lys	Asp 215	Arg	Asn	Leu	Gly	Phe 220	Lys	Ile	Ser	Leu
His 225	Суз	Pro	Суз	Суз	Thr 230	Phe	Val	Pro	Ser	Asn 235	Asn	Tyr	Ile	Ile	Pro 240
Asn	Lys	Ser	Glu	Glu 245	Leu	Glu	Ala	Arg	Phe 250	Ala	Gly	Ile	Asp	Gly 255	Thr
Ser	Thr	Tyr	Thr 260	Ser	Gly	Asp	Gln	Lys 265	Thr	Ile	Lys	Ser	Thr 270	Arg	Lys
Lys	Asn	Ser 275	Gly	Lys	Thr	Pro	His 280	Leu	Leu	Leu	Met	Leu 285	Leu	Pro	Ser
Tyr	Arg 290	Leu	Glu	Ser	Gln	Gln 295	Thr	Asn	Arg	Arg	300	Lys	Arg	Ala	Leu
Asp 305	Ala	Ala	Tyr	Cys	Phe 310	Arg	Asn	Val	Gln	Asp 315	Asn	Cys	Cys	Leu	Arg 320
Pro	Leu	Tyr	Ile	Asp 325	Phe	Lys	Arg	Asp	Leu 330	Gly	Trp	Lys	Trp	Ile 335	His
Glu	Pro	Lys	Gly 340	Tyr	Asn	Ala	Asn	Phe 345	Cys	Ala	Gly	Ala	Cys 350	Pro	Tyr
Leu	Trp	Ser 355	Ser	Asp	Thr	Gln	His 360	Ser	Arg	Val	Leu	Ser 365	Leu	Tyr	Asn
Thr	Ile 370	Asn	Pro	Glu	Ala	Ser 375	Ala	Ser	Pro	Суз	Суз 380	Val	Ser	Gln	Asp
Leu 385	Glu	Pro	Leu	Thr	Ile 390	Leu	Tyr	Tyr	Ile	Gly 395	Lys	Thr	Pro	Lys	Ile 400

Glu Gln Leu Ser Asn Met Ile Val Lys Ser Cys Lys Cys Ser

405 410

<210> 4

<211> 412

<212> PRT

<213> Homo sapien

<400> 4

Met Lys Met His Leu Gln Arg Ala Leu Val Val Leu Ala Leu Leu Asn $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Phe Ala Thr Val Ser Leu Ser Leu Ser Thr Cys Thr Thr Leu Asp Phe 20 25 30

Gly His Ile Lys Lys Lys Arg Val Glu Ala Ile Arg Gly Gln Ile Leu $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Ser Lys Leu Arg Leu Thr Ser Pro Pro Glu Pro Thr Val Met Thr His 50 55 60

Val Pro Tyr Gln Val Leu Ala Leu Tyr Asn Ser Thr Arg Glu Leu Leu 65 70 75 80

Glu Glu Met His Gly Glu Arg Glu Glu Gly Cys Thr Gln Glu Asn Thr
85 90 95

Glu Ser Glu Tyr Tyr Ala Lys Glu Ile His Lys Phe Asp Met Ile Gln 100 105 110

Gly Leu Ala Glu His Asn Glu Leu Ala Val Cys Pro Lys Gly Ile Thr 115 120 125

Ser Lys Val Phe Arg Phe Asn Val Ser Ser Val Glu Lys Asn Arg Thr 130 135 140

Ser Lys Arg Asn Glu Gln Arg Ile Glu Leu Phe Gln Ile Leu Arg Pro 165 170 175

Asp Glu His Ile Ala Lys Gln Arg Tyr Ile Gly Gly Lys Asn Leu Pro 180 185 190

Thr	Arg	Gly 195	Thr	Ala	Glu	Trp	Leu 200	Ser	Phe	Asp	Val	Thr 205	Asp	Thr	Val
Arg	Glu 210	Trp	Leu	Leu	Arg	Arg 215	Glu	Ser	Asn	Leu	Gly 220	Leu	Glu	Ile	Ser
Ile 225	His	Суз	Pro	Суз	His 230	Thr	Phe	Gln	Pro	Asn 235	Gly	Asp	Ile	Leu	Glu 240
Asn	Ile	His	Glu	Val 245	Met	Glu	Ile	Lys	Phe 250	Lys	Gly	Val	Asp	Asn 255	Glu
Asp	Asp	His	Gly 260	Arg	Gly	Asp	Leu	Gly 265	Arg	Leu	Lys	Lys	Gln 270	Lys	Asp
His	His	Asn 275	Pro	His	Leu	Ile	Leu 280	Met	Met	Ile	Pro	Pro 285	His	Arg	Leu
Asp	Asn 290	Pro	Gly	Gln	Gly	Gly 295	Gln	Arg	Lys	Lys	Arg 300	Ala	Leu	Asp	Thr
Asn 305	Tyr	Cys	Phe	Arg	Asn 310	Leu	Glu	Glu	Asn	Cys 315	Cys	Val	Arg	Pro	120
Tyr	Ile	Asp	Phe	Arg 325	Gln	Asp	Leu	Gly	Trp 330	Lys	Trp	Val	His	Glu 335	Pro
Lys	Gly	Tyr	Tyr 340	Ala	Asn	Phe	Cys	Ser 345	Gly	Pro	Cys	Pro	Tyr 350	Leu	Arg
Ser	Ala	Asp 355	Thr	Thr	His	Ser	Thr 360	Val	Leu	Gly	Leu	Tyr 365	Asn	Thr	Leu
Asn	Pro 370	Glu	Ala	Ser	Ala	Ser 375	Pro	Cys	Cys	Val	Pro 380	Gln	Asp	Leu	Glu
Pro 385	Leu	Thr	Ile	Leu	Tyr 390	Tyr	Val	Gly	Arg	Thr 395	Pro	Lys	Val	Glu	Gln 400
Leu	Ser	Asn	Met	Val	Val	Lys	Ser	Cys	Lys	Cys	Ser				

```
<210> 5
<211> 18
<212> PRT
<213> rat
<400> 5
Met Lys Trp Val Thr Phe Leu Leu Leu Phe Ile Ser Gly Ser Ala
                                 10
Phe Ser
<210> 6
<211> 36
<212> DNA
<213> Chinese hamster
<400> 6
gcgatacccg ggtataccat ggccacctca gcaagt
                                                                   36
<210> 7
<211> 33
<212> DNA
<213> Chinese hamster
<400> 7
cgggtgttcg aattagtttt tgtattggaa ggg
                                                                   33
<210> 8
<211> 56
<212> DNA
<213> Homo sapien
<400> 8
                                                                   56
ggttctgcct tttctcacca ccatcaccac caccatcatc tgtccacctg caagac
<210> 9
<211> 30
<212> DNA
<213> Homo sapien
<400> 9
tagtctcgag ttatcagctg cacttgcagg
                                                                   30
<210> 10
<211> 45
<212> DNA
<213> rat
```

<400> 10

```
45
aaagggggat ccgccaccat gaagtgggta acctttctcc tcctc
<210> 11
<211> 45
<212> DNA
<213> rat
<400> 11
agaaaaggca gaaccggaga tgaagaggag gaggagaaag gttac
                                                                    45
<210> 12
<211> 39
<212> DNA
<213> pig
<400> 12
                                                                    39
cgcctcagtg cccactgttc ctgtgacagc aaagataac
<210> 13
<211> 39
<212> DNA
<213> pig
<400> 13
                                                                    39
gttatctttg ctgtcacagg aacagtgggc actgaggcg
<210> 14
<211> 33
<212> DNA
<213> pig
<400> 14
                                                                    33
ggatccctgt ccacctccaa gaccatcgac atg
<210> 15
<211> 33
<212> DNA
<213> pig
<400> 15
                                                                    33
catgtcgatg gtcttggagg tggacaggga tcc
<210> 16
<211> 20
<212> PRT
<213> synthetic
<400> 16
His His His His His His Leu Ser Thr Ser Lys Thr Ile Asp
```

10

15

```
Met Glu Leu Val
20
```

<210> 17 <211> 20 <212> PRT <213> pig

<400> 17

Ala Leu Asp Thr Asn Tyr Cys Phe Ser Ser Thr Glu Lys Asn Cys Cys 1 5 10 15

Val Arg Gln Leu 20